

FIG. 1

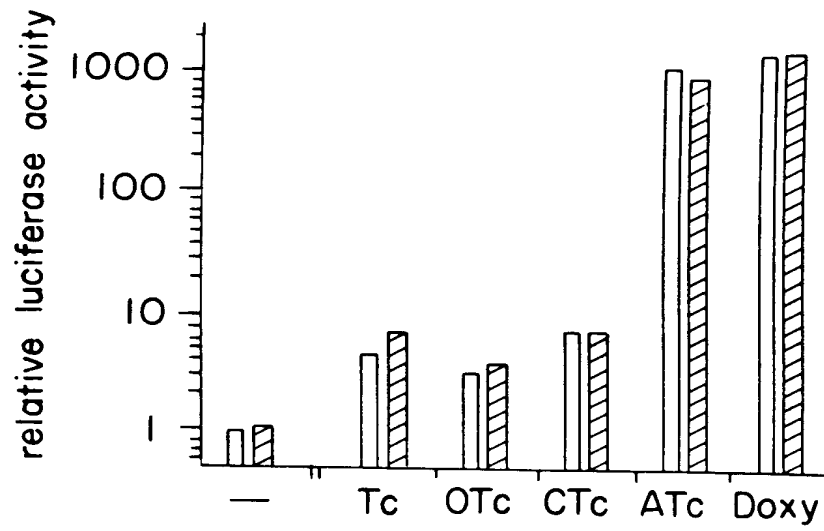


FIG.2

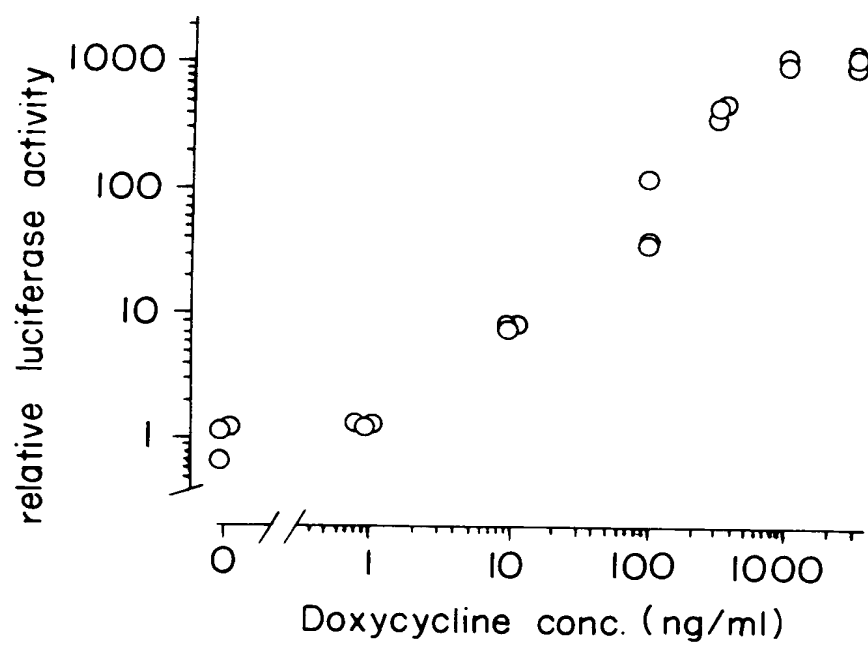
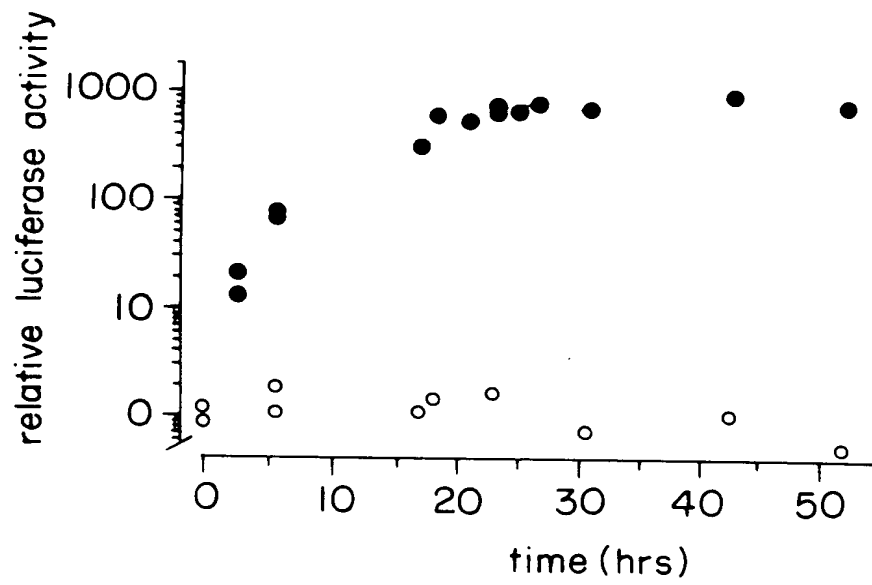


FIG. 3



H T H

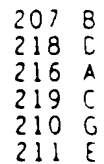
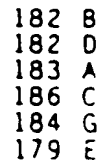
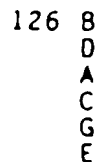
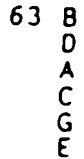


FIG. 5

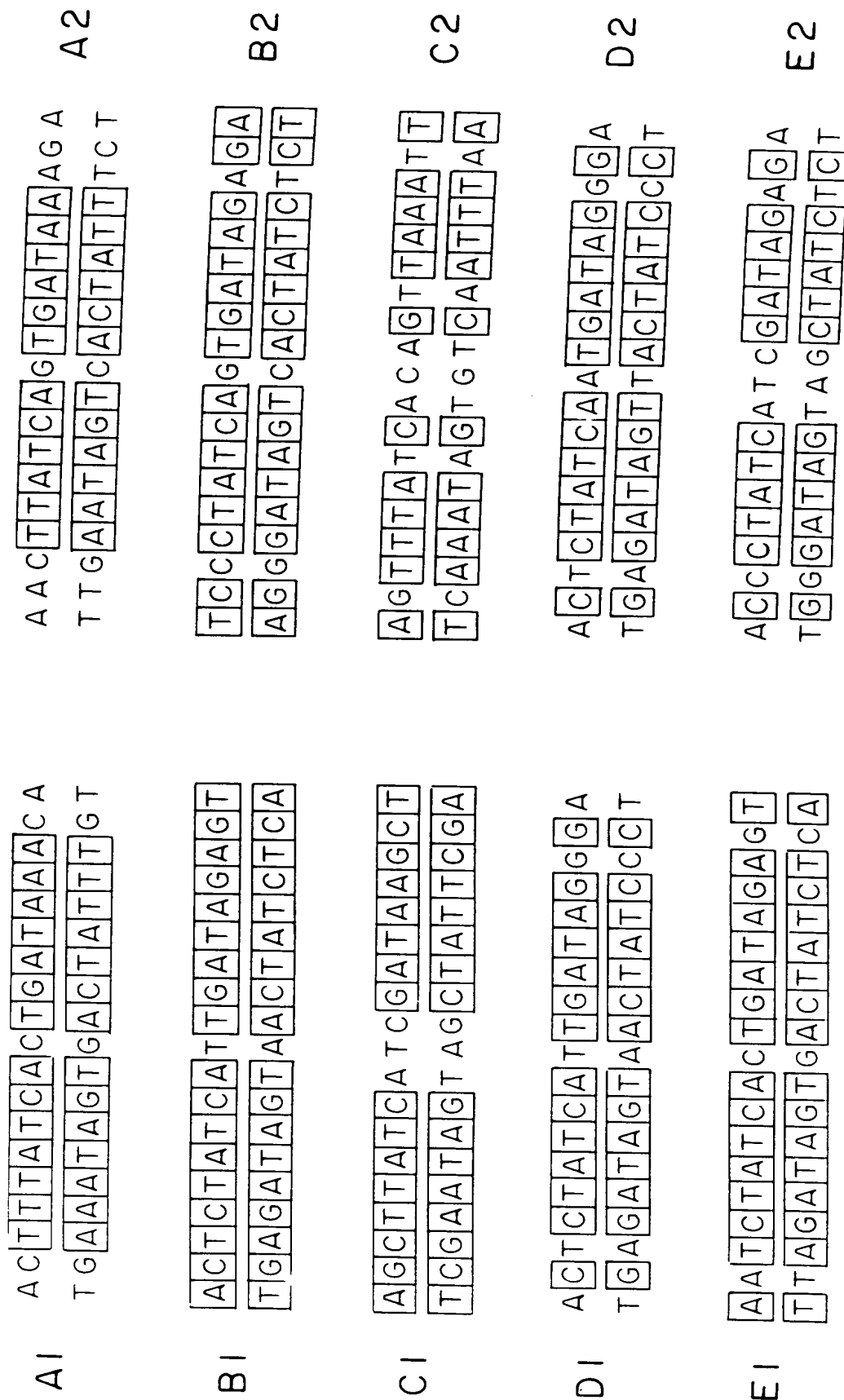


FIG. 6

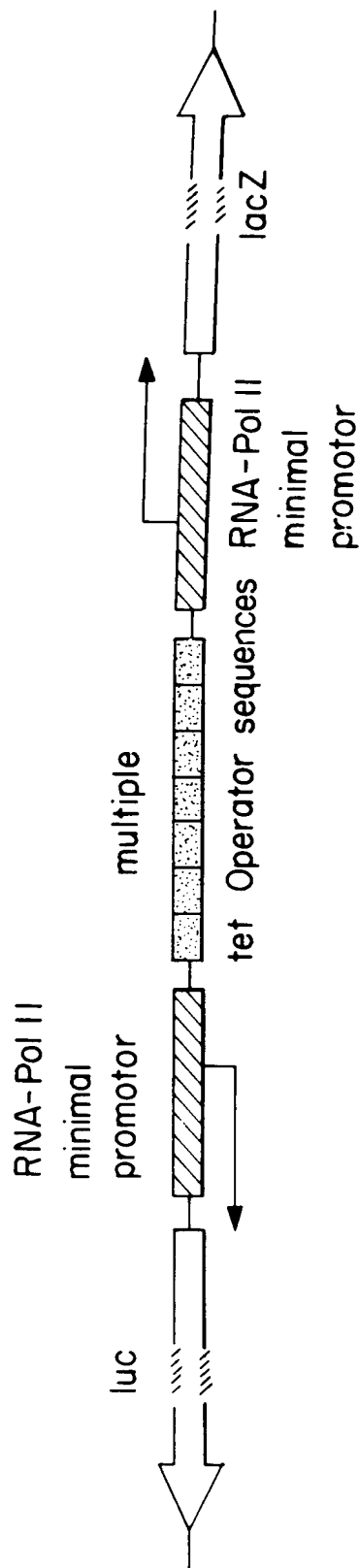


FIG. 7A

5' GAATTCGGGG
EcoRI + 75

CCGCGGAGGCTGGATCGGTCCCGGTGTCTTCTATGGAGGTCAAAACAGCGTGGA

+ 1
←
C

TGGCGTCTCCAGGCGATCTGACGGTTCCTAAACGAGCTCTGCTTATATAGG
P_{hCMV}*-3 -3;

tet O

TC (GAGTTTACCACTCCCTATCAGTGATAGAGAAAAGTGAAAGTC)₇GAGC

TCGGTACCCGGGTCGAGTAGGCGTGTACGGTGGGAGGCCTATATAAGCAGAG
P_{hCMV}*-1
-53

CTCGTTTAGTGAACCGTCAGATCGCCTGGAGACGCCATCCACGCTGTTTTGA
+ 1 →

CCTCCATAGAAGACACCGGGACCGATCCAGCCTCCGCGGCCCCCGAATTC 3'
+ 75 EcoRI

FIG. 7B

5' ⁺¹⁹ AGATCT GCAGGGTCGC
Bgl II Pst I

← ⁺¹
A
TCGGTGTTCGAGGCCACACGCGTCACCTT AATATGCGAAGTGGACC GGATC
P_{K*} -37 -37

tet O
TC (GAGTTTACCACTCCCTATC AGTGATAGAGAAAAGTGAAAGTC)₇ GAGC

TCGGTACCCGGGTCGAGTAGGCGTGTACGGTGGGAGGCCTATATAAGCAGAG
-53 P_{hCMV*} -1

CTCGTTTAGTGAACCGTCAGATCGCCTGGAGACGCCATCCACGCTGTTTTGA
+1 →

CCTCCATAGAAGACACCGGGACCGATCCAGCCTCCGCGGCCCCG GAATTC 3'
+75 EcoRI

000000 24E 4460

FIG. 8A

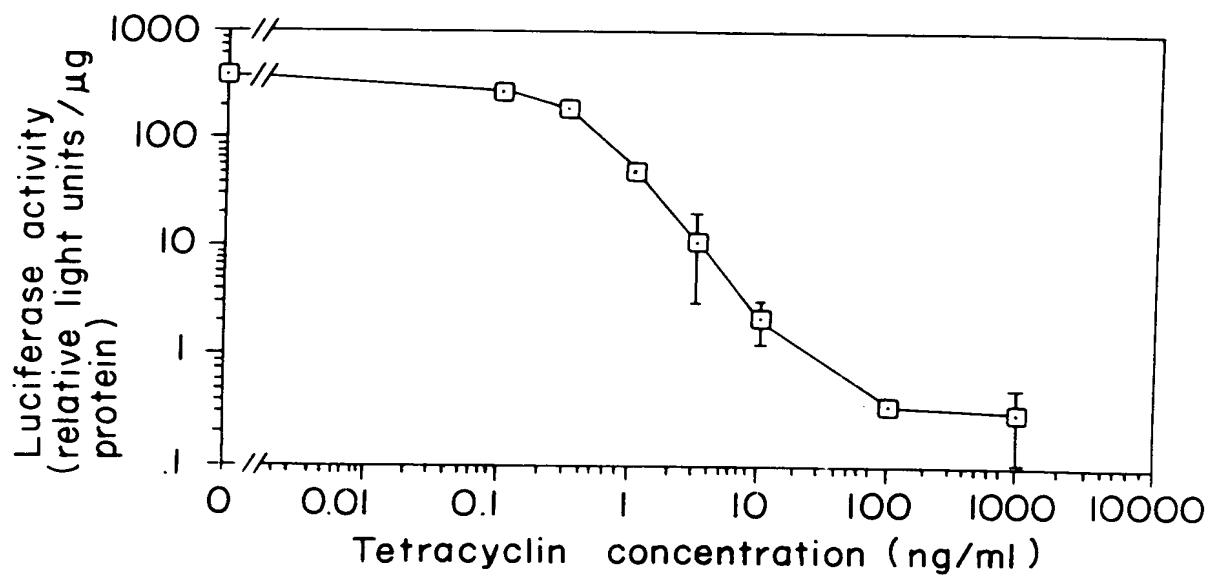


FIG. 8B

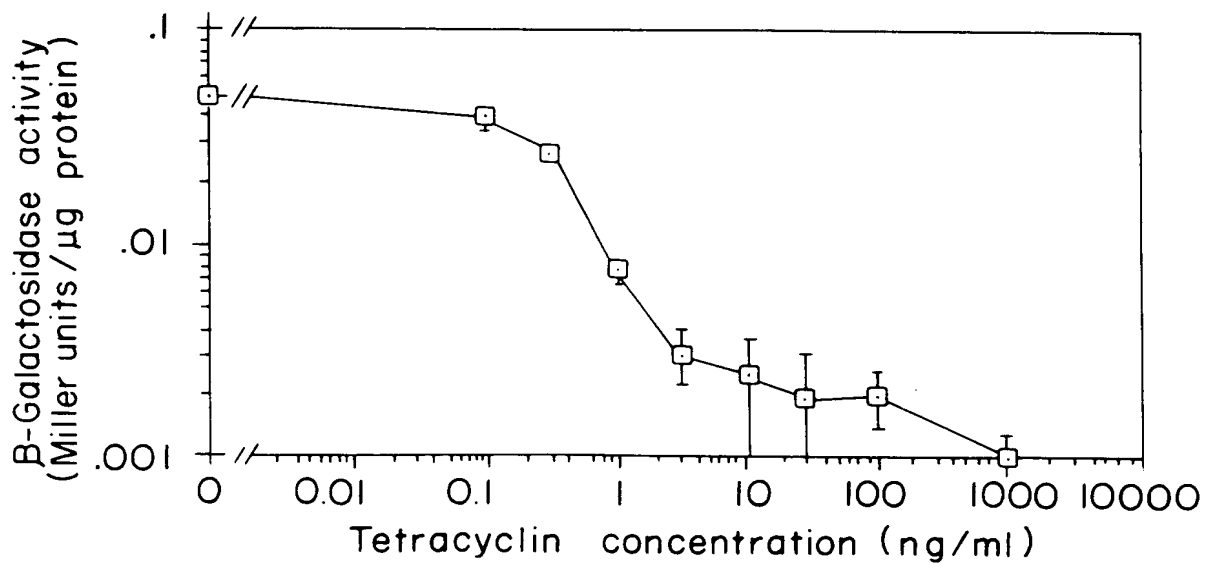


FIG. 9A

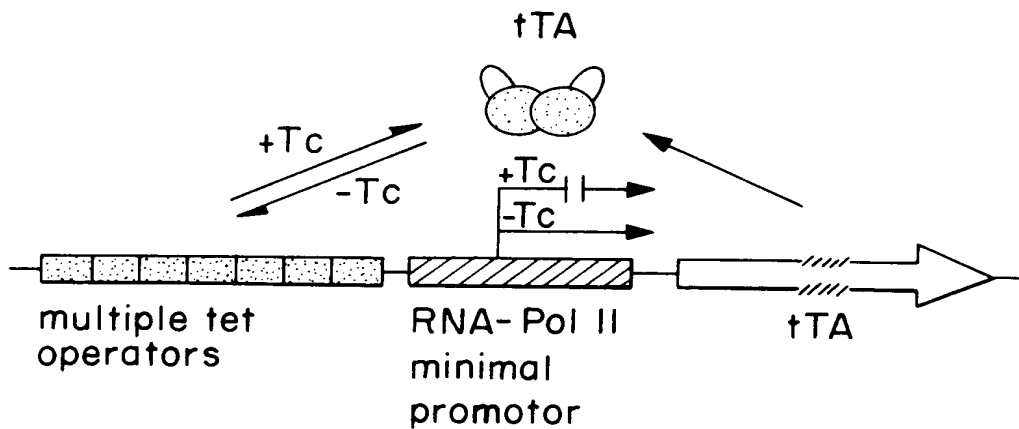


FIG. 9B

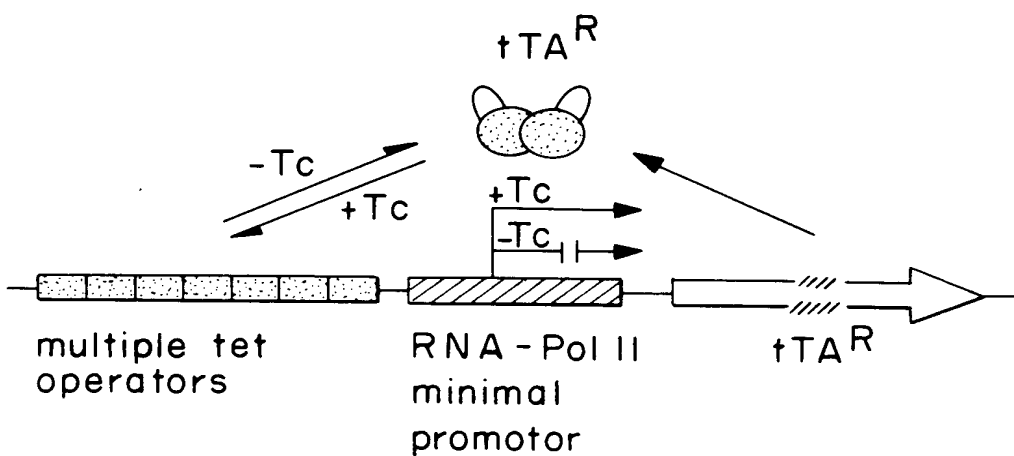


FIG.10

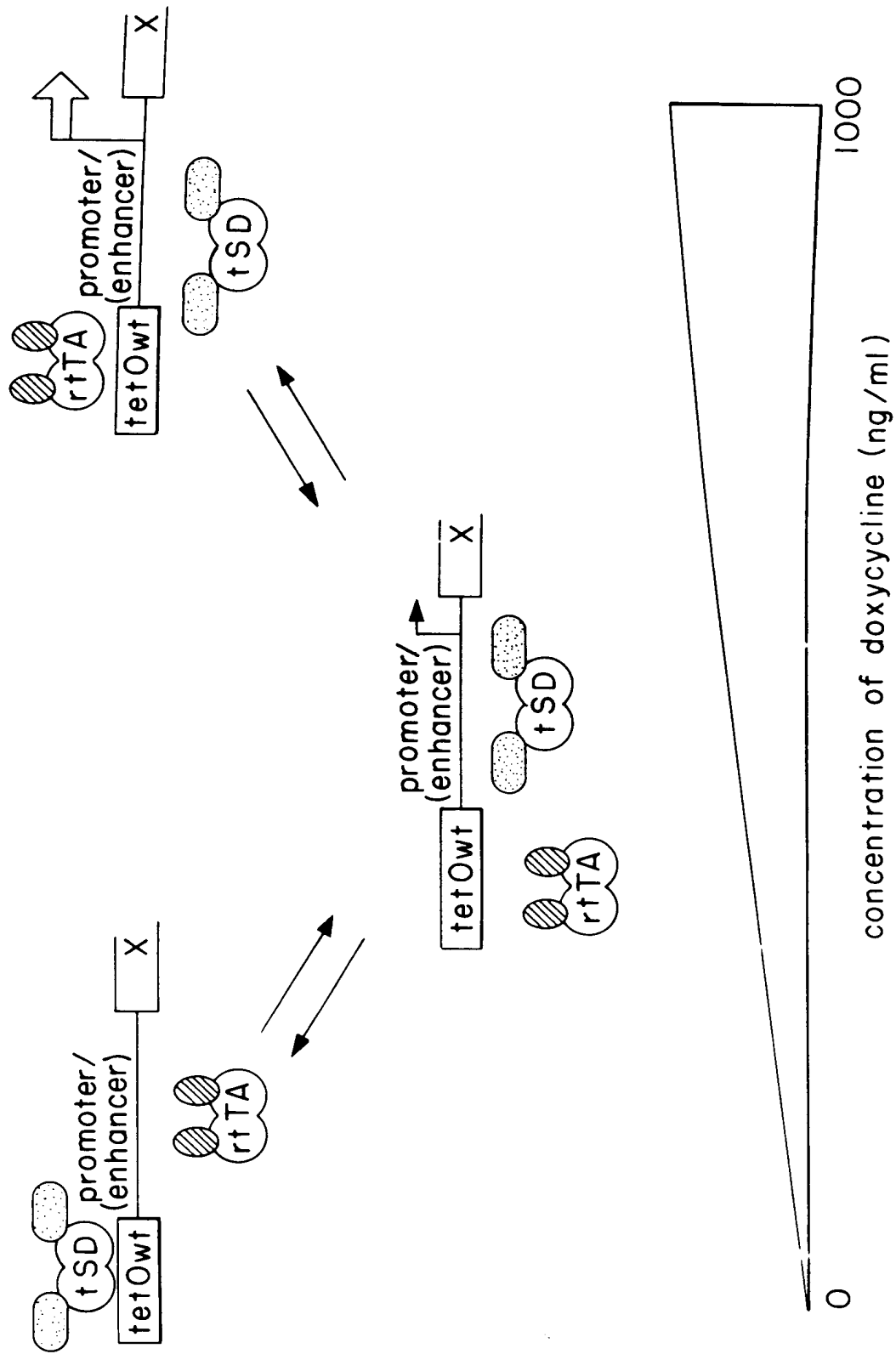


FIG. 11

pUHD141Kr-1

Pcmv →

pUHD141sma-1

ATG TCT... tetR gene ... TCC CCG GGT AAC TAAGTAAGGATCC — An —

in-frame fusion of the Krueppel repr./silencing domain

pUHD141erb-1

in-frame fusion of the v-erbA repr./silencing domain

FIG.12

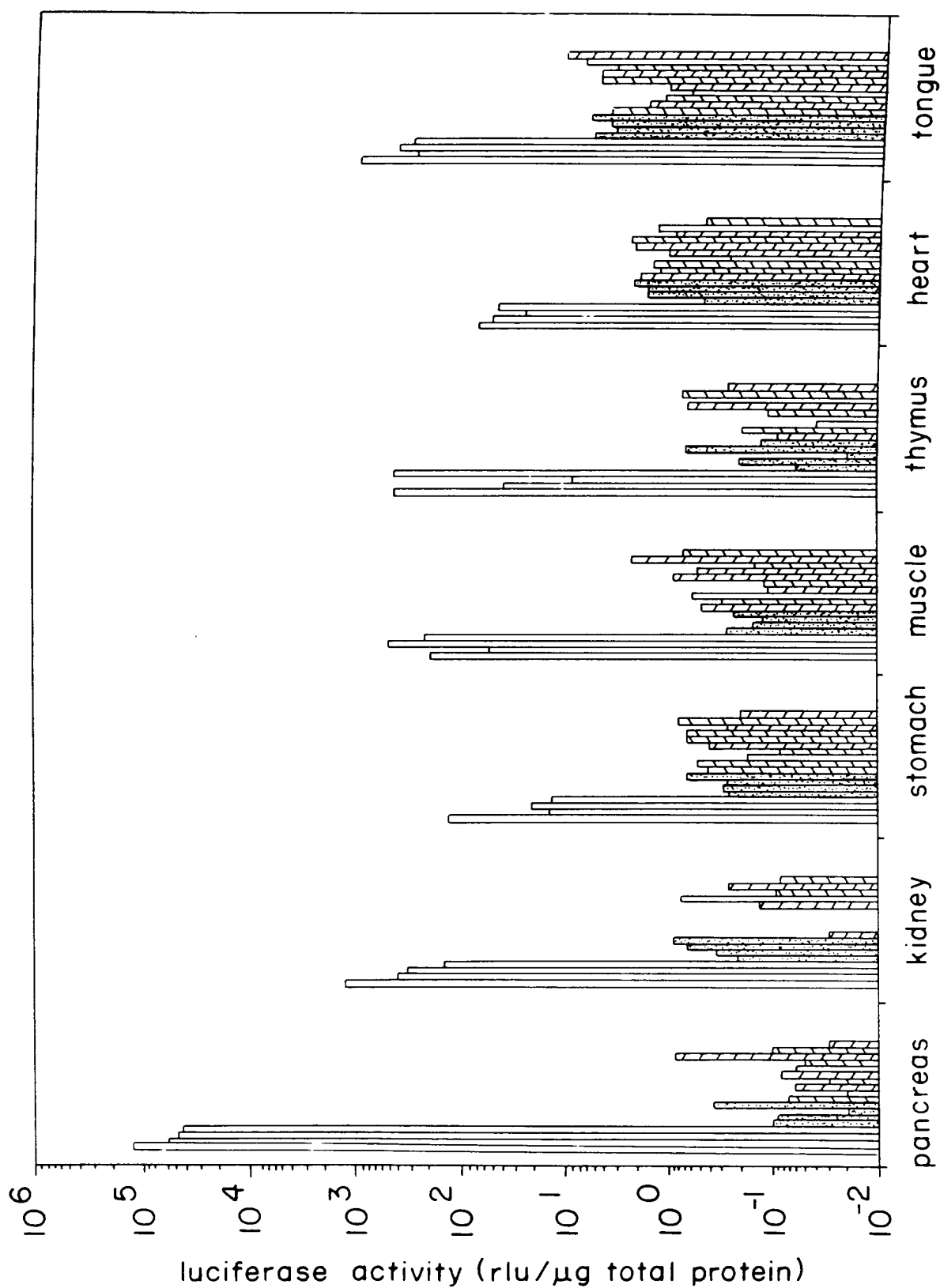


FIG. 13

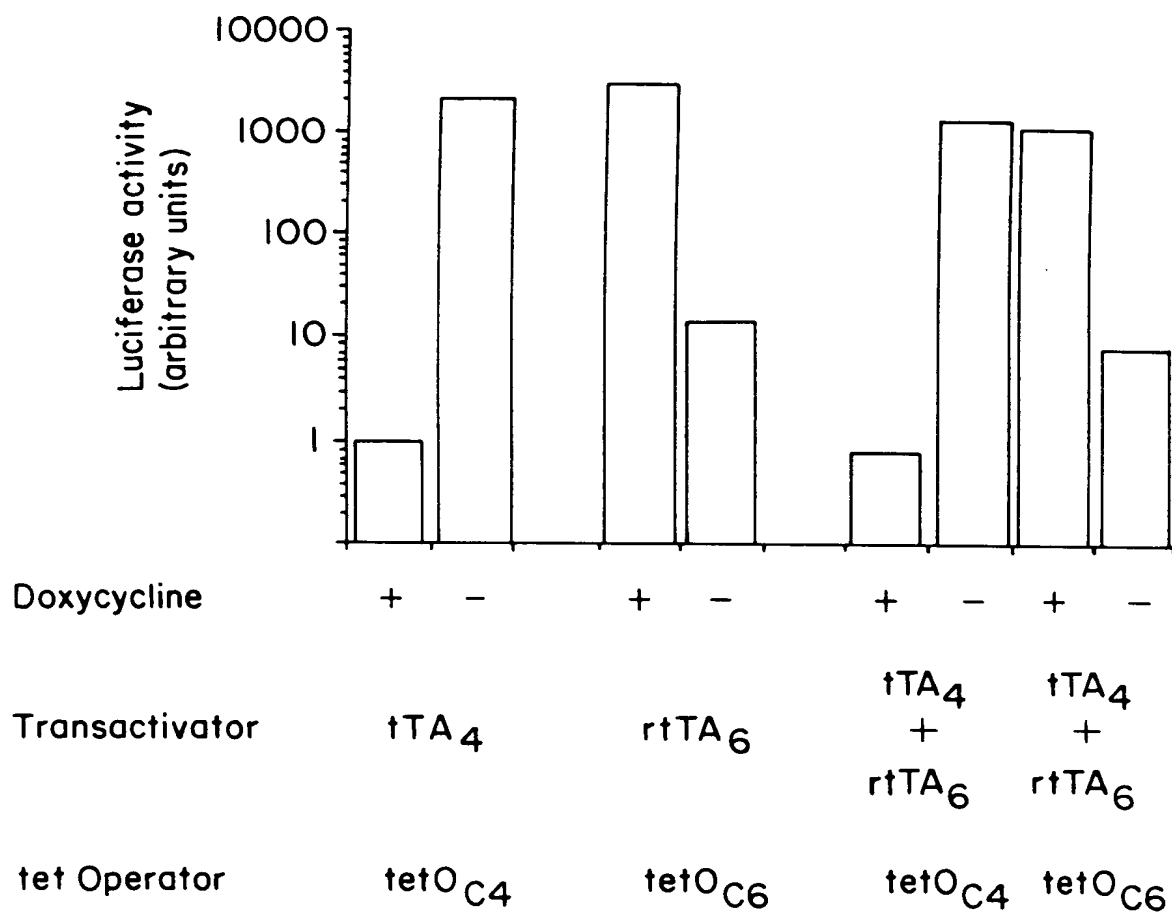


FIG.14A

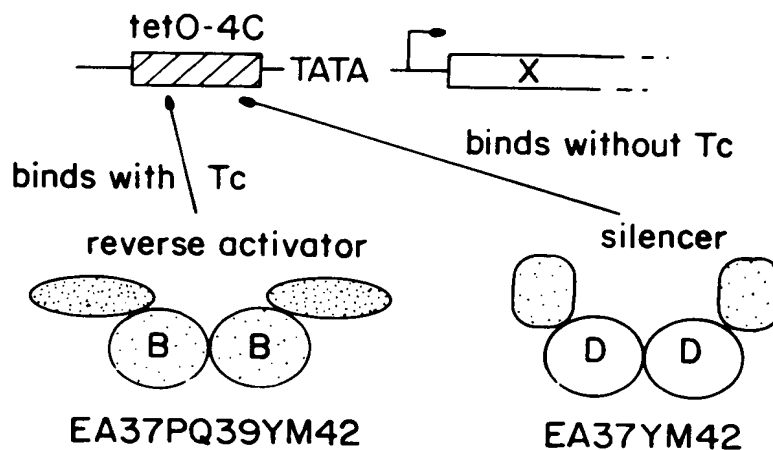


FIG.14B

